

### SYR-HSD-5002 Substitute Sequence Listing.ST25.txt SEQUENCE LISTING

SYRRX, INC.

<120> PROBE, ASSAY AND KITS FOR DETECTING 11B-HYDROXYSTEROID DEHYDROGENASE AND MODULATORS THEREOF

<130> SYR-HSD-5002-U

us 10/800,140 <140>

2004-03-11 <141>

<160> 7

<170> PatentIn version 3.2

<210>

292 <211>

<212> **PRT** 

<213> Homo sapiens

<220>

MISC\_FEATURE <221>

<222> (1)..(292)

<223> Amino acid sequence for full-length human wild type 11B-hydroxysteroid dehydrogenase

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Met Ala Phe Met Lys Lys Tyr Leu Leu Pro Ile Leu Gly Leu Phe Met 1 10 15

Ala Tyr Tyr Tyr Ser Ala Asn Glu Glu Phe Arg Pro Glu Met Leu 20 25 30

Gln Gly Lys Lys Val Ile Val Thr Gly Ala Ser Lys Gly Ile Gly Arg
35 40 45

Glu Met Ala Tyr His Leu Ala Lys Met Gly Ala His Val Val Thr 50 60

Ala Arg Ser Lys Glu Thr Leu Gln Lys Val Val Ser His Cys Leu Glu 65 70 75 80

Leu Gly Ala Ala Ser Ala His Tyr Ile Ala Gly Thr Met Glu Asp Met 85 90 95

Thr Phe Ala Glu Gln Phe Val Ala Gln Ala Gly Lys Leu Met Gly Gly

Leu Asp Met Leu Ile Leu Asn His Ile Thr Asn Thr Ser Leu Asn Leu 115 120

Phe His Asp Asp Ile His His Val Arg Lys Ser Met Glu Val Asn Phe Page 1

Leu Ser Tyr Val Val Leu Thr Val Ala Ala Leu Pro Met Leu Lys Gln 145 150 155 160

Ser Asn Gly Ser Ile Val Val Ser Ser Leu Ala Gly Lys Val Ala 165 170 175

Tyr Pro Met Val Ala Ala Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly 180 185 190

Phe Phe Ser Ser Ile Arg Lys Glu Tyr Ser Val Ser Arg Val Asn Val 195 200 205

Ser Ile Thr Leu Cys Val Leu Gly Leu Ile Asp Thr Glu Thr Ala Met 210 215 220

Lys Ala Val Ser Gly Ile Val His Met Gln Ala Ala Pro Lys Glu Glu 225 230 235 240

Cys Ala Leu Glu Ile Ile Lys Gly Gly Ala Leu Arg Gln Glu Glu Val 245 250 255

Tyr Tyr Asp Ser Ser Leu Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys 260 265 270

Arg Lys Ile Leu Glu Phe Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg 275 280 285

Phe Ile Asn Lys 290

<210> 2

<211> 286

<212> PRT <213> Artificial

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<223> Amino acid sequence for residues 24-292 of 11B-hydroxysteroid dehydrogenase with a N-terminal MKHQHQHQHQHQQPL tag

<400> 2

Met Lys His Gln His Gln His Gln His Gln His Gln Gln Pro
5 10 15

Leu Asn Glu Glu Phe Arg Pro Glu Met Leu Gln Gly Lys Lys Val Ile 20 25 30

Val Thr Gly Ala Ser Lys Gly Ile Gly Arg Glu Met Ala Tyr His Leu Page 2 Ala Lys Met Gly Ala His Val Val Val Thr Ala Arg Ser Lys Glu Thr 50 60 Leu Gln Lys Val Val Ser His Cys Leu Glu Leu Gly Ala Ala Ser Ala 65 70 75 80 His Tyr Ile Ala Gly Thr Met Glu Asp Met Thr Phe Ala Glu Gln Phe 85 90 95 Val Ala Gln Ala Gly Lys Leu Met Gly Gly Leu Asp Met Leu Ile Leu 100 105 110 Asn His Ile Thr Asn Thr Ser Leu Asn Leu Phe His Asp Asp Ile His 115 120 125 His Val Arg Lys Ser Met Glu Val Asn Phe Leu Ser Tyr Val Val Leu 130 135 140 Thr Val Ala Ala Leu Pro Met Leu Lys Gln Ser Asn Gly Ser Ile Val 145 150 160 Val Val Ser Ser Leu Ala Gly Lys Val Ala Tyr Pro Met Val Ala Ala 165 170 175 Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly Phe Phe Ser Ser Ile Arg 180 185 190 Lys Glu Tyr Ser Val Ser Arg Val Asn Val Ser Ile Thr Leu Cys Val 195 200 205 Leu Gly Leu Ile Asp Thr Glu Thr Ala Met Lys Ala Val Ser Gly Ile 210 215 220 Val His Met Gln Ala Ala Pro Lys Glu Glu Cys Ala Leu Glu Ile Ile 225 230 235 240 Lys Gly Gly Ala Leu Arg Gln Glu Glu Val Tyr Tyr Asp Ser Ser Leu 245 250 255 Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys Arg Lys Ile Leu Glu Phe 260 265 270 Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg Phe Ile Asn Lys 275 280 285

<210> 3 <211> 1405 <212> DNA

<213> Artificial

<220>

<223> Human cDNA sequence encoding residues 24-292 of 11B-hydroxysteroid dehydrogenase

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<sup>&</sup>lt;210> 4 <211> 24 <212> DNA <213> Artificial

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<210> <211> <212> <213>	6 23 DNA Artificial	
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<400> gatctte	7 cctg gatggatttc tga	23

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<110> SYRRX, INC. <120> PROBE, ASSAY AND KITS FOR DETECTING 11B-HYDROXYSTEROID DEHYDROGENASE AND MODULATORS THEREOF <130> SYR-HSD-5002-U us 10/800,140 <140> <141> 2004-03-11 <160> 7 <170> PatentIn version 3.2 <210> <211> 292 <212> **PRT** <213> Homo sapiens <220> <221> MISC\_FEATURE <222> (1)..(292)Amino acid sequence for full-length human wild type 11B-hydroxysteroid dehydrogenase <400> 1 Met Ala Phe Met Lys Lys Tyr Leu Leu Pro Ile Leu Gly Leu Phe Met  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ Ala Tyr Tyr Tyr Ser Ala Asn Glu Glu Phe Arg Pro Glu Met Leu 20 25 30 Gln Gly Lys Lys Val Ile Val Thr Gly Ala Ser Lys Gly Ile Gly Arg 35 40 45 Glu Met Ala Tyr His Leu Ala Lys Met Gly Ala His Val Val Thr 50 60 Ala Arg Ser Lys Glu Thr Leu Gln Lys Val Val Ser His Cys Leu Glu 65 70 75 80 Leu Gly Ala Ala Ser Ala His Tyr Ile Ala Gly Thr Met Glu Asp Met
85 90 95 Thr Phe Ala Glu Gln Phe Val Ala Gln Ala Gly Lys Leu Met Gly Gly Leu Asp Met Leu Ile Leu Asn His Ile Thr Asn Thr Ser Leu Asn Leu 115 120 125

Phe His Asp Asp Ile His His Val Arg Lys Ser Met Glu Val Asn Phe Page 1 Leu Ser Tyr Val Val Leu Thr Val Ala Ala Leu Pro Met Leu Lys Gln 145 150 155 160

Ser Asn Gly Ser Ile Val Val Val Ser Ser Leu Ala Gly Lys Val Ala 165 170 175

Tyr Pro Met Val Ala Ala Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly
180 185 190

Phe Phe Ser Ser Ile Arg Lys Glu Tyr Ser Val Ser Arg Val Asn Val 195 200 205

Ser Ile Thr Leu Cys Val Leu Gly Leu Ile Asp Thr Glu Thr Ala Met 210 220

Lys Ala Val Ser Gly Ile Val His Met Gln Ala Ala Pro Lys Glu Glu 225 230 235 240

Cys Ala Leu Glu Ile Ile Lys Gly Gly Ala Leu Arg Gln Glu Glu Val 245 250 255

Tyr Tyr Asp Ser Ser Leu Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys 260 265 270

Arg Lys Ile Leu Glu Phe Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg 275 280 285

Phe Ile Asn Lys 290

<210> 2

<211> 286

<212> PRT <213> Artificial

<220>

<223> Amino acid sequence for residues 24-292 of 11B-hydroxysteroid dehydrogenase with a N-terminal MKHQHQHQHQHQHQPL tag

<400> 2

Met Lys His Gln His Gln His Gln His Gln His Gln His Gln Pro
1 5 10 15

Leu Asn Glu Glu Phe Arg Pro Glu Met Leu Gln Gly Lys Lys Val Ile 20 25 30

Val Thr Gly Ala Ser Lys Gly Ile Gly Arg Glu Met Ala Tyr His Leu Page 2 Ala Lys Met Gly Ala His Val Val Thr Ala Arg Ser Lys Glu Thr 50 60

Leu Gln Lys Val Val Ser His Cys Leu Glu Leu Gly Ala Ala Ser Ala 65 70 75 80

His Tyr Ile Ala Gly Thr Met Glu Asp Met Thr Phe Ala Glu Gln Phe 85 90 95

Val Ala Gln Ala Gly Lys Leu Met Gly Gly Leu Asp Met Leu Ile Leu 100 105 110

Asn His Ile Thr Asn Thr Ser Leu Asn Leu Phe His Asp Asp Ile His 115 120 125

His Val Arg Lys Ser Met Glu Val Asn Phe Leu Ser Tyr Val Val Leu 130 135 140

Thr Val Ala Ala Leu Pro Met Leu Lys Gln Ser Asn Gly Ser Ile Val 145 150 155 160

Val Val Ser Ser Leu Ala Gly Lys Val Ala Tyr Pro Met Val Ala Ala 165 170 175

Tyr Ser Ala Ser Lys Phe Ala Leu Asp Gly Phe Phe Ser Ser Ile Arg 180 185 190

Lys Glu Tyr Ser Val Ser Arg Val Asn Val Ser Ile Thr Leu Cys Val 195 200 205

Leu Gly Leu Ile Asp Thr Glu Thr Ala Met Lys Ala Val Ser Gly Ile 210 215 220

Val His Met Gln Ala Ala Pro Lys Glu Glu Cys Ala Leu Glu Ile Ile 225 230 235 240

Lys Gly Gly Ala Leu Arg Gln Glu Glu Val Tyr Tyr Asp Ser Ser Leu 245 250 255

Trp Thr Thr Leu Leu Ile Arg Asn Pro Cys Arg Lys Ile Leu Glu Phe 260 265 270

Leu Tyr Ser Thr Ser Tyr Asn Met Asp Arg Phe Ile Asn Lys 275 280 285

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